

```

1 1 ACA GTC AGC CGC ATG GCT CCC CTG TGC CCC AGC CCC TGG CTC CTG CCT CTG L P L 12
1 1 ACA GTC AGC CGC ATG GCT CCC CTG TGC CCC AGC CCC TGG CTC CTG CCT CTG L P L 48

13 1 L I P A P A P G L T V Q L L L S 28
49 TTG ATC CCG GCC CCT GCT CCA GGC CTC ACT GTG CAA CTG CTG CTG TCA 96

29 1 L L L M P V H P Q R L P R M Q 44
97 CTG CTG CTT CTG ATG CCT GTC CAT CCC CAG AGG TTG CCC CGG ATG CAG 144

45 1 E D S P L G G G S S G E D D P L 60
145 GAG GAT TCC CCC TTG GGA GGA GGC TCT TCT GGG GAA GAT GAC CCA CTG 192

61 1 G E E D L P S E E D S P R E E D 76
193 GGC GAG GAG GAT CTG CCC AGT GAA GAG GAT TCA CCC AGA GAG GAG GAT 240

77 1 P P G E E D L P G E E D L P G E 92
241 CCA CCC GGA GAG GAG GAT CTA CCT GGA GAG GAT CTA CCT GGA GAG 288

93 1 E D L P E V K P K S E E E G S L 108
289 GAG GAT CTA CCT GAA GTT AAG CCT AAA TCA GAA GAA GAG GGC TCC CTG 336

109 1 K L E D L P T V E A P G D P Q E 124
337 AAG TTA GAG GAT CTA CCT ACT GTT GAG GCT CCT GGA GAT CCA GAA 384

125 1 P Q N N A H R D K E G D D Q S H 140
385 CCC CAG AAT AAT GCC CAC AGG GAC AAA GAA GGG GAT GAC CAG AGT CAT 432

141 1 W R Y G G D P P W P R V S P A C 156
433 TGG CGC TAT GGA GGC GAC CCG CCC TGG CCC CGG GTG TCC CCA GCC TGC 480

157 1 A G R F Q S P V D I R P Q L A A 172
481 GCG GGC CGC TTC CAG TCC CCG CCG GTG GAT ATC CGC CCC CAG CTC GCC GCC 528

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FIG.-1A

173 F C P A L R P L F E G L F E G G F Q L P 188
 529 TTC TGC CCG GCC CTG CGC CCC CTG GAA CTC CTG GGC TTC CAG CTC CCG 576

 189 P L P E L R L R L R N N G H S V Q L 204
 577 CCG CTC CCA GAA CTG CGC CTG CGC AAC AAT GGC CAC AGT GTG CAA CTG 624

 205 T L P P G L E M A L G P G R E Y 220
 625 ACC CTG CCT CCT GGG CTA GAG ATG GCT CTG GGT CCC GGG CGG GAG TAC 672

 221 R A L Q L H L H W G A A G R P G 236
 673 CGG GCT CTG CAG CTG CAT CTG CAC TGG GGG GCT GCA GGT CGT CCG GGC 720

 237 S E H T V E G H R F P A E I H V 252
 721 TCG GAG CAC ACT GTG GAA GGC CAC CGT TTC CCT GCC GAG ATC CAC GTG 768

 253 V H L S T A F A R V D E A L G R 268
 769 GTT CAC CTC AGC ACC GCC TTT GCC AGA GTT GAC GAG GCC TTG GGG CGC 816

 269 P G G L A V L A A F L E E G P E 284
 817 CCG GGA GGC CTG GCC GTG TTG GCC TTT CTG GAG GAG GGC CCG GAA 864

 285 E N S A Y E Q L L S R L E E I A 300
 865 GAA AAC AGT GCC TAT GAG CAG CTG TTG TCT CGC TTG GAA GAA ATC GCT 912

 301 E E G S E T Q V P G L D I S A L 316
 913 GAG GAA GGC TCA GAG ACT CAG GTC CCA GGA CTG GAC ATA TCT GCA CTC 960

 317 L P S D F S R Y F Q Y E G S L T 332
 961 CTG CCC TCT GAC TTC AGC CGC TAC TTC CAA TAT GAG GGG TCT CTG ACT 1008

 333 T P P C A Q G V I W T V F N Q T 348
 1009 ACA CCG CCC TGT GCC CAG GGT GTC ATC TGG ACT GTG TTT AAC CAG ACA 1056

349 V M L S A K Q L T T H E G T A A L E S S E D D T L W 364
 1057 GTG ATG CTG AGT GCT AAG CAG CTC CAC ACC CTC TCT GAC ACC CTG TGG 1104
 365 G P G D S R L Q L N F R A T Q P 380
 1105 GGA CCT GGT GAC TCT CGG CTA CAG CTG AAC TTC CGA GCG ACG CAG CCT 1152
 381 L N G R V I E A S F P A G V D S 396
 1153 TTG AAT GGG CGA GTG ATG ATT GAG GCC TTC CCT GCT GGA GTG GAC AGC 1200
 397 S P R A A E P V Q L N S C L A A 412
 1201 AGT CCT CGG GCT GCT GAG CCA GTC CAG CTG AAT TCC TGC CTG GCT GCT 1248
 413 G D I L A L V F G L L F A V T S 428
 1249 GGT GAC ATC CTA GCC CTG GTT TTT GGC CTC CTT TTT GCT GTC ACC AGC 1296
 429 V A F L V Q M R R Q H R R G T K 444
 1297 GTC GCG TTC CTT GTG CAG ATG AGA AGG CAG CAC AGA AGG GGA ACC AAA 1344
 445 G G V S Y R P A E V A E T G A * 460
 1345 GGG GGT GTG AGC TAC CGC CCA GCA GAG GTA GCC GAG ACT GGA GCC TAG 1392
 1393 AGG CTG GAT CTT GGA GAA TGT GAG AAG CCA GCC AGA GGC ATC TGA GGG 1440
 1441 GGA GCC GGT AAC TGT CCT GTC CTG CTC ATT ATG CCA CTT CCT TTT AAC 1488
 1489 TGC CAA GAA ATT TTT TAA AAT AAA TAT TTA TAA T 1522

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FIG._1C

FIG._1

FIG._1A

FIG._1B

FIG._1C

1 ggatcctgtt gactcgtgac cttaccccc accctgtgct ctctgaaaca tgagctgtgt
 61 ccactcaggg ttaaatggat taaggcggtt gcaagatgtg ctttgtaaa cagatgcttg
 121 aaggcagcat gctcgttaag agtcatcacc aatccctaata ctcaagtaata cagggacaca
 181 aacactgcgg aaggccgcag ggtcctctgc ctaggaaaaa cagagacctt tggtcacttg
 241 ttatctgac cttccctcca ctattgtcca tgaccctgcc aatccccctt ctgtgagaaa
 301 caccacaaga ttatcaataa aaaaataaat ttaaaaaaaa aatacaaaaa aaaaaaaa
 361 aaaaaaaa gacttacgaa tagttattga taaatgaata gctattggtg aagccaagta
 421 aatgatcata ttcaaaaacca gacggccatc atcacagctc aagtctacct gatttgatct
 481 ctttatcatt gtcattcttt ggattcacta gattagtcac catcctcaaa attctcccc
 541 aagttctaata tacgttccaa acatttaggg gttacatgaa gcttgaacct actaccttct
 601 ttgcttttga gccatgagtt gtaggaatga tgagtttaca cctacatgc tggggattaa
 661 tttaaaacttt accttaagt cagttgggtg gcctttggct tatttttcta cctccacct
 721 tagttaatgg atgcactgtg ttacagtaatt gcttacctaa gacctaaag cctatttctc
 781 gggtaggtag gtactcagtt ttacagtaatt gcttacctaa gacctaaag cctatttctc
 841 ttgtactggc ctttatctgt aatatgggca tatttaatac aataaattt ttggagtttt
 901 ttgtgtgtt tggtgtgtt ttgtgtgtt acggagtctt gcatctgtca tggccaggct
 961 ggagtagcag tgggtccatc tcggctcact gcaagctcca cctccgagt tcacgccatt
 1021 ttcctgcctc agcctccga gtagctggga ctacaggcg cggccacct gccgggctaa
 1081 ttttttgtat ttttggtaga gacggggttt caccgtgtta gccagaatgg tctcgatctc
 1141 ctgacttcgt gatccacccg cctcggcctc ccaaagtctt gggattacag gtgtgagcca
 1201 ccgcacctgg ccaattttt gagtctttaa aagtaaaaat atgtcttcta agctggtaac
 1261 tatggtaaat ttccttttat taatgtggtg ctgacgggtca tatagggttct tttaggtttg
 1321 gcatgcataat gctactttt gcagtccttt cattacattt ttctctctc atttgaagag
 1381 catgttatat ctttttagctt cacttggctt aaaagggttct ctcattagcc taacacagt
 1441 tcattgttgg taccacttgg atcataagt gaaaaacagt caagaaatg cacagtaata
 1501 cttgtttgta agagggatga ttcagggtgaa tctgacacta agaaactccc ctacctgagg
 1561 tctgagattc ccttgacatt gctgtatata ggcttttctt ttgacagcct gtgactgcgg
 1621 actattttcc ttaagcaaga tatgctaaaag ttttgtgagc ctttttccag agagaggtct
 1681 catatctgca tcaagtgaga acatataatg tctgcatgtt tccatatctc aggaatgttt
 1741 gcttgtgttt tatgctttta tatagacagg gaaacttgtt cctcagtgac ccaaaagagg
 1801 tgggaattgt tattggatat catcattggc ccacgcttct tgaccttggg aacaattaa
 1861 ggttcataat ctcaattctg tcagaattgg tacaagaaat agctgctatg ttcttgaca
 1921 tccacttgg taggaaataa gaatgtgaaa ctcttcagtt ggtgtgtgtc cct?gtttt

FIG.-2A

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1981 ttgcaatttc cttcttactg tgttaaaaa aagtatgac ttgctctgag agtgaggca
2041 ttcttaatca tgatctttaa agatcaataa tataatcctt tcaaggatta tgtctttatt
2101 ataataaaga taatttgtct ttaacagaat caataatata atcccttaa ggattatc
2161 ttgtctgggc gcagtggctc acacctgtaa tcccagcact ttgggtggcc aagtggaag
2221 gatcaaatctt gctacttctt atattatctt ctaaagcaga attcatctct ctccctcaa
2281 tatgatgata ttgatgggt ttgccctcac tcactagatt gtgagctcct gctcaggga
2341 ggtagcgttt ttgttttttg tttttgtttt tcttttttga gacaggtctt tgctctgtca
2401 ccaggccag agtgcaatgg tacagtctca gctcactgca gcctcaaccg cctcggctca
2461 aaccatcatc ccatttcagc ctccctgagta gctgggacta caggcacatg ccattacac
2521 tggctaattt ttttgtattt ctagttaga cagggtttgg ccatgttgcc cgggctggtc
2581 tcgaactcct ggactcaagc aatccaccca cctcagcctc ccaaaatgag ggaccgtgtc
2641 ttattcatct ccattgtcct agtccatagc ccagtgtggt acctatggta gtactaaata
2701 aatatttgtt gaatgcaata gtaaatagca tttcaggag caagaactag attaacaag
2761 gtggtaaaag gtttgagaa aaaaataata gtttaatttg gctagagtat gaggagagt
2821 agtaggagac aagatggaaa ggtctcttgg gcaaggtttt gaaggaagt ggaagtcaga
2881 agtacacaat gtgcatactg tggcaggcag tggggagcca atgaaggctt ttgagcagga
2941 gagtaattgt ttgaaaaata aatataggtt aaacctata gagccctctt gacacataca
3001 cttgcttttc attcaagctc aagtttgtct ccacatacc cattacttaa ctcacctcg
3061 ggctccctta gcagcctgcc ctacctctt acctgcttcc tgggtggagt agggatgtat
3121 acatgagctg cttccctct cagccagagg acatgggggg cccagctcc cctgccttc
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3301 ccattggccc gataaccttc tgcctgtgca cacacctgcc cctcactcca ccccatcct
3361 agcttttgta tgggggagag ggcacagggc cagacaaaac tgtgagactt tggctccatc
3421 tctgcaaaag ggcgctctgt gagtcaagcct gctccctcc aggcttgctc ctccccacc
3481 cagctctcgt ttccaatgca cgtacagccc gtacacaccg tgtgctggga caccctACAG
3541 TCAGCCGCAT GGCTCCCTG TGCCCCAGCC CTTGGCTCCC TCTGTTGATC CCGGCCCTG
3601 CTCCAGGCCT CACTGTGCAA CTGTGCTGT CACTGTGCTT TCTGGTGCCT GTCCATCCCC
3661 AGAGGTTGCC CCGGATGCAG GAGGATTCCC CCTTGGGAGG AGGCTCTTCT GGGGAAGATG
3721 ACCCACTGGG CGAGGAGGAT CTGCCCAGTG AAGAGGATTC ACCCAGAGAG GAGGATCCAC
3781 CCGGAGAGGA GGATCTACCT GGAGAGGAGG ATCTACCTGG AGAGGAGGAT CTACCTGAAG
3841 TTAAGCCCTAA ATCAGAAGAA GAGGGCTCCC TGAAGTTAGA GGATCTACCT ACTGTTGAGG
3901 CTCCCTGGAGA TCCTCAAGAA CCCCAGAATA ATGCCCACAG GGACAAAGAA Ggtaagtgg

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FIG. 2C

5941 CGGGAGGCCT GGCGGTGTG GCCGCTTTC TGGAGGtacc agatcctgga caccacctac
 6001 tccccgcttt ccatcccat gctcctccc gactctatcg tggagccaga gaccacctcc
 6061 cagcaagctc actcaggccc ctggctgaca aactcattca cgcactgttt gttcatttaa
 6121 caccactgt gaaccaggca ccagccccc acaaggattc tgaagctgta ggtccttgcc
 6181 tctaaggagc ccacagccag tgggggaggg tgacatgaca gacacatagg aaggacatag
 6241 taaagatggg ggtcacagag gagggtgacac ttaaagcctt cactggtaga aaagaaaagg
 6301 aggtgttcat tgcagaggaa acagaatgtg caaagactca gaatatggcc tatttaggga
 6361 atggctacat acaccatgat tagaggaggc ccagtaaagg gaagggatgg tgagatgcct
 6421 gctaggttca ctactcact ttattttatt tattttatt ttgacagtc tctctgtcgc
 6481 ccaggctgga gtgcagtgg gtgatcttgg gtcactgcaa ctccgcctc ccgggttcaa
 6541 gggattctcc tgcctcagct tctgagtag ctggggttac aggtgtgtgc caccatgccc
 6601 agctaatttt ttttgtatt tttagtagac aggttttcac catgttggtc aggtggtct
 6661 caaactcctg gcctcaagt gctcgcctga ctcagcctac caaagtgtg attacaagt
 6721 tgagccaccg tgcccagcca cactcactga ttctttaatg ccagccacac agcacaaagt
 6781 tcagagaaat gcctccatca tagcatgtca atatgttcat actcttaggt tcatgatgtt
 6841 cttaacatta ggttcataag caaataaaga aaaagaata ataaataaaa gaagtggcat
 6901 gtcaggacct cacctgaaa gccaaacaca gaatcatgaa ggtgaatgca gaggtgacac
 6961 caacacaaag gtgtatatat ggtttcctgt ggggagtatg tacggaggca gcagtgagt
 7021 agactgcaaa cgtcagaagg gcacgggtca ctgagagcct agtatcctag taaagtggc
 7081 tctctccctc tctctccagc ttgtcattga aaaccagtc accaagcttg ttggttcgca
 7141 cagcaagagt acatagagt tgaataata cataggatt taagaggag acactgtctc
 7201 taaaaaaa aacaacagca acaacaaaa gcaacaacca ttacaatttt atgttccctc
 7261 agcatcttca gagctgagga atgggagagg actatgggaa ccccttcat gttccggcct
 7321 tcagccatgg ccctggatac atgcaactcat ctgtcttaca atgtcattcc ccagGAGGG
 7381 CCCGGAAGAA AACAGTGCCT ATGAGCAGTT GCTGTCTCGC TTGGAAGAAA TCGCTGAGGA
 7441 AGgtcagttt gttggtctgg ccaataatct ctgtggccta gttcataaag aatcacctt
 7501 tggagcttca ggtctgaggc tggagatggg ctccctccag tgcaggaggg attgaagcat
 7561 gagccagcgc tcatcttgat aataaccatg aagctgacag acacagttac ccgcaaacgg
 7621 ctgcctacag attgaaaaac aagcaaaaac cgccgggcac ggtggctcac gcctgtaac
 7681 ccagcacttt gggaggccaa ggcagggtgga tcacgaggtc aagagatcaa gaccatcctg
 7741 gccaacatgg tgaaaaccca tctctactaa aaatacgaaa aaatagccag gctggtggc
 7801 ggtgctctgt aatcccagct actcgggagg ctgaggcagg agaattggcat gaaccggga
 7861 ggcagaagtt gcagtgagcc gagatcgtgc cactgcactc cagcctgggc aacagagcga

FIG._2D

7921 gactcttgct tcaaaaaaa aaaaaaaa gaaacccaag caaaaaccaa aatgagacaa
 7981 aaaaaaag accaaaaat ggtgtttgga aattgtcaag gtcaagtctg gagagctaaa
 8041 ctttttctga gaactgttta tctttaataa gcatcaataa ttttaacttt gtaaatactt
 8101 ttgttggaat tctgttctct tagtcact cttgggtcat tttaaatctc acttactcta
 8161 ctgaccttt ctgaccttt taggtttctg ctgacctgt tagaactctg ctttgcatt tcttgtgtct
 8221 gttttgtata gttatcaata ttatattta ttacaagt ttacagatca ttttttctt
 8281 tctttttttt tttttttttt ttttttttat ctgacctgt gtttagtaga gacagggttt caccatattg
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 8461 atggtagaca gagttaagag ttagactca gacgggtctt cttcttctct tctcttctct
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 8641 agggcctgca cttagtgag aagtgtctc agagttgagt taccttggt tctgggaggt
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 8761 tagatcctct tcacagGCTC AGAGACTCAG GTCCACAGGAC TGGACATATC TGCACCTCTG
 8821 CCTCTGACT TCAGCCGCTA CTTCCAATAT GAGGGTCTC TGACTACACC GCCCTGTGCC
 8881 CAGGGTGCTA TCTGGACTGT GTTTAACCCAG ACAGTGATGC TGAGTGCTAA GCAGgtgggc
 8941 ctgggtgtg tgtggacaca gtgggtgctg gggaaagag atgtaagatg agatgagaaa
 9001 caggagaaga aagaaatcaa ggctgggctc tgtggcttac gcctataatc ccaccagtt
 9061 gggaggctga ggtggagaa tggttttagc ccaggagttc aagacaaggc ggggcaacat
 9121 agtgtgacct catctctacc tactcaagga ggctgaggtg ggaagatcgc ttgattccag
 9181 gtagcggcc tagtccagc ctatgatccc accactgcct accatcttta ggatacattt
 9241 gagtttgaga ctgcagttag aagaggctgg atggggaata caggagctgg agggaggagc
 9301 atttatttat cctgaggtgc cactgacct ccttagCTCC ACACCTCTC TGACACCTG TGGGGACCTG
 9361 cctgaggtgc cctgaggtgc cactgacct ccttagCTCC ACACCTCTC TGACACCTG TGGGGACCTG
 9421 ccacactgt GTGACTCTCG GCTACAGCTG AACTTCCGAG CGACGCAGCC TTTGAATGGG CGAGTGATTG
 9481 GTGACTCTCG GCTACAGCTG AACTTCCGAG CGACGCAGCC TTTGAATGGG CGAGTGATTG
 9541 AGGCTCTCTT CCCTGCTGGA GTGGACAGCA GTCCTCGGGC TGCTGAGCCA Ggtacagctt
 9601 tgtctgggtt ccccccagcc agtagtccc tatcctccc tatcctccc tgtgtgtgct agtgtctgtc
 9661 attggtggtc acagcccgcc tctcacatct cctttttctc tccagTCCAG CTGAATTCTT
 9721 GCCTGGCTGC TGgtgagttc gcccctctc ttggtcctga tggcaggaga ctcctcagca
 9781 ccattcagcc ccagggtgc tcaggaccgc ctctgctccc tctcctttc tgcagaacag
 9841 accccaacc caatataga gaggcagatc atggtgggga tcccccat tccccagag

9901 gctaattgat tagaatgaag cttgagaaat ctccagcat ccctctcgca aaagaatccc
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 10141 ttggctttta ggaagcaaaa acggtgctta tcttaccct tctcgtgat ccaccctcat
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 10261 ggggtggctg agtgcaactga ggcagggtgtt gaggaactct gcagaccctt ctctctccc
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 10561 gAAGGGGAAC CAAAGGGGT GTGAGCTACC GCCCAGCAGA GTAGCCGAG ACTGGAGCCT
 10621 AGAGGCTGGA TCTTGGAGAA TGTGAGAAC CAGCCAGAGG CATCTGAGGG GGAGCCGGTA
 10681 ACTGTCCTGT CCTGCTCATT ATGCCACTTC CTTTAACTG CCAAGAAATT TTTTAAATA
 10741 AATATTATA Aataaatatg tgtagtcac ctttgttccc caaatcagaa ggaggtattt
 10801 gaatttccta ttactgttat tagcaccaat ttagtggttaa tgcattttatt ctattacagt
 10861 tcggcctcct tccacacatc actccaatgt gttgctcc

FIG._2F

FIG. 2A

FIG. 2B

FIG. 2C

FIG. 2D

FIG. 2E

FIG. 2F

FIG._2

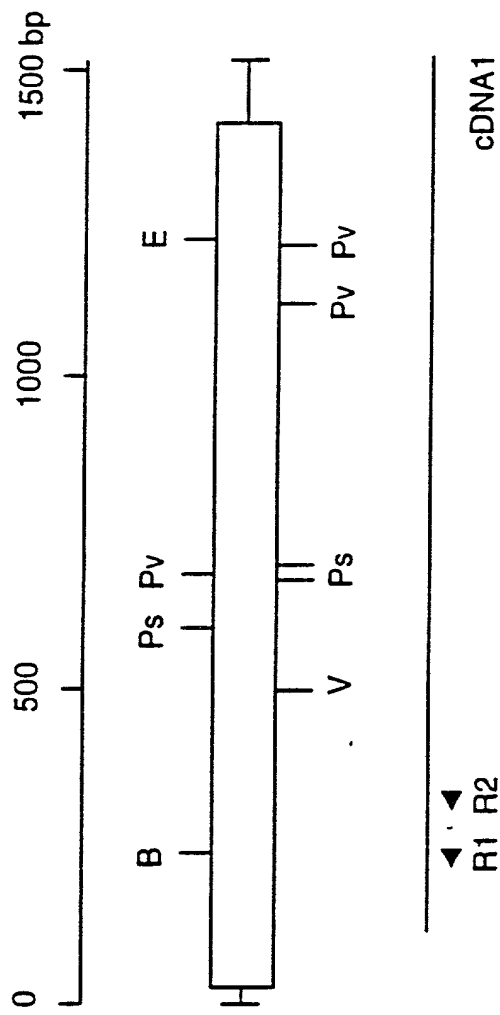


FIG. 3

5' MN Genomic Region

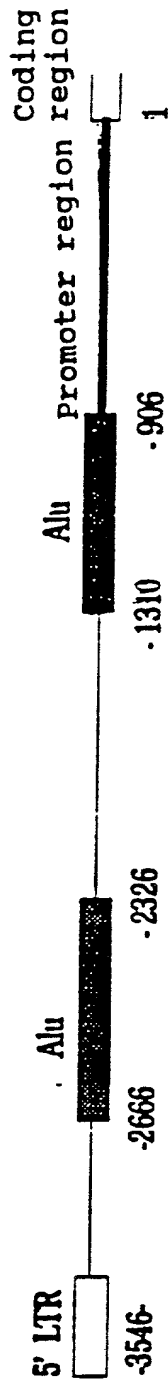


FIG._4

-506 CTTGCTTTTC ATTCAAGCTC AAGTTGTCT CCCACATACC CATTACTAA CTCACCCCTCG

-446 GGCTCCCCCTA GCAGCCTGCC CTACCTCTTT ACCTGCTTCC TGGTGGAGTC AGGGATGTAT
AP2

-386 ACATGAGCTG CTTTCCCTCT CAGCCAGAGG ACATGGGGGG CCCACAGCTCC CCTGCCCTTC

-326 CCCTTCTGTG CCTGGAGCTG GGAAGCAGGC CAGGGTTAGC TGAGGCTGGC TGGCAAGCAG

-266 CTGGGTGGTG CCAGGGAGAG CCTGCATAGT GCCAGGTGGT GCCTTGGGT CCAAGCTAGT
VII p53

-206 CCATGGCCCC GATAACCTTC TGCCTGTGCA CACACCTGCC CCTCACTCCA CCCCCATCCT
VI Inr V

-146 AGCTTTGGTA TGGGGGAGAG GGCACAGGC CAGACAAACC TGTGAGACTT TGGCTCCATC
IV AP1 III Inr

-86 TCTGCAAAAG GCGCTCTGT GAGTCAGCCT GCTCCCCCTCC AGGCTTGCTC CTCCCCCACC
II AP1 p53 I AP2

-26 CAGCTCTCGT TTCCAATGCA CGTACAGCCC GTACACACCG TGTGCTGGGA CACCCACAG

FIG._6

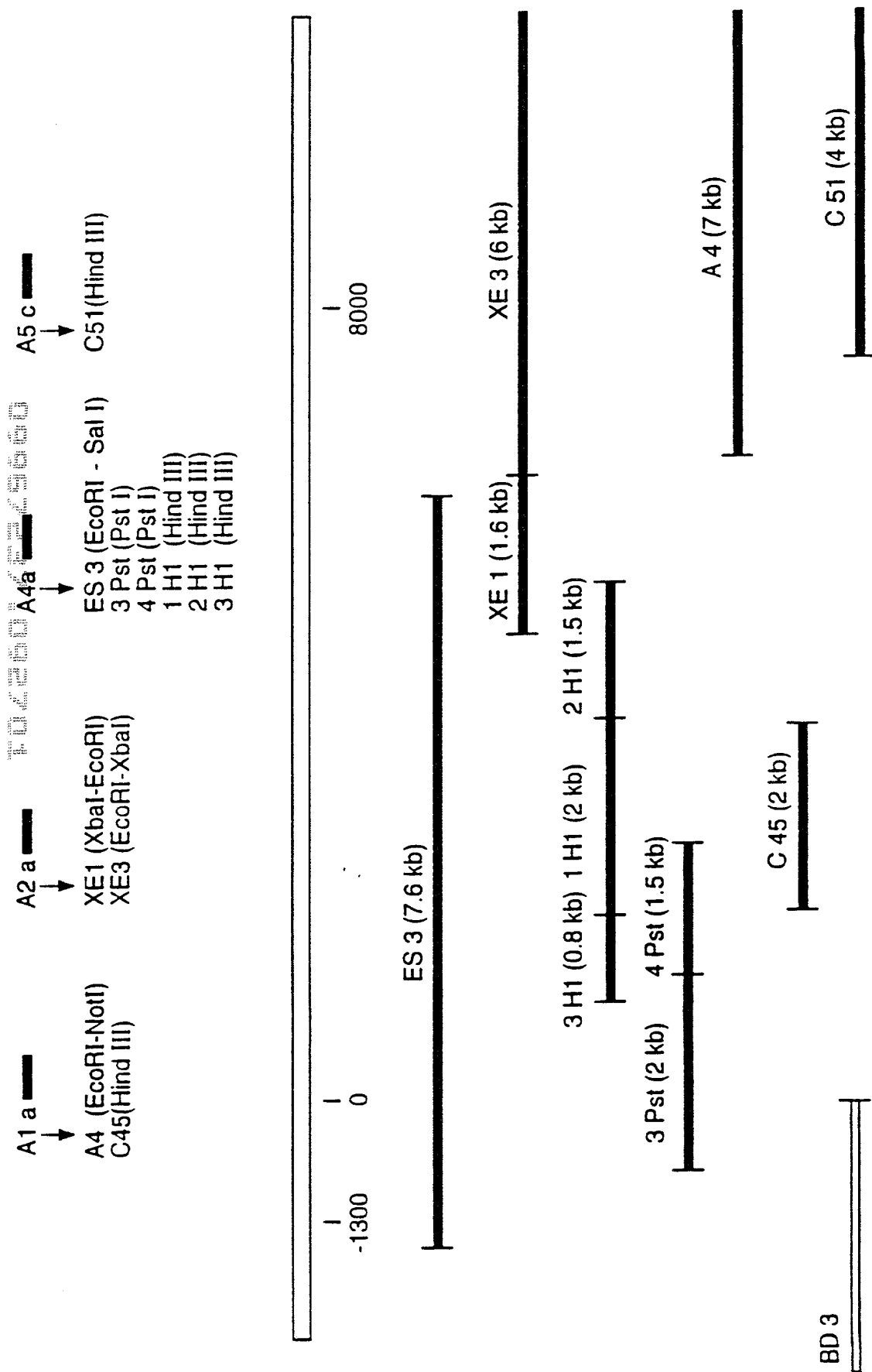


FIG. 7

1kb

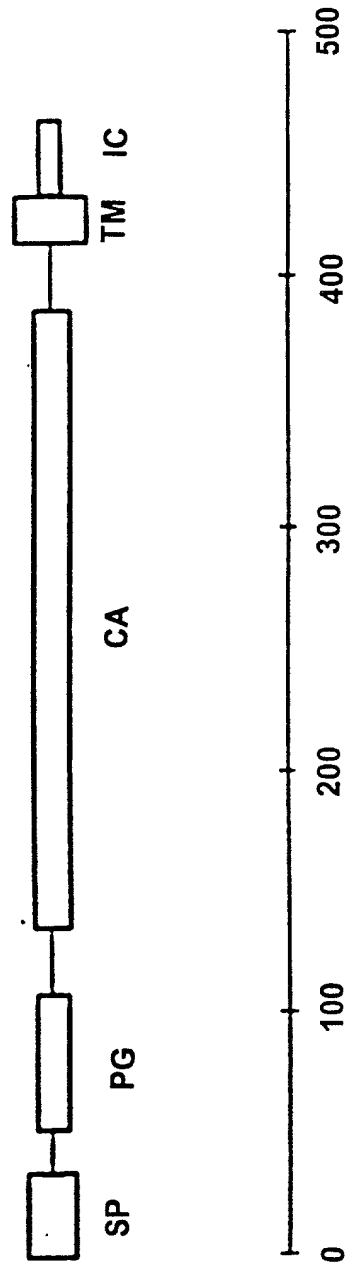
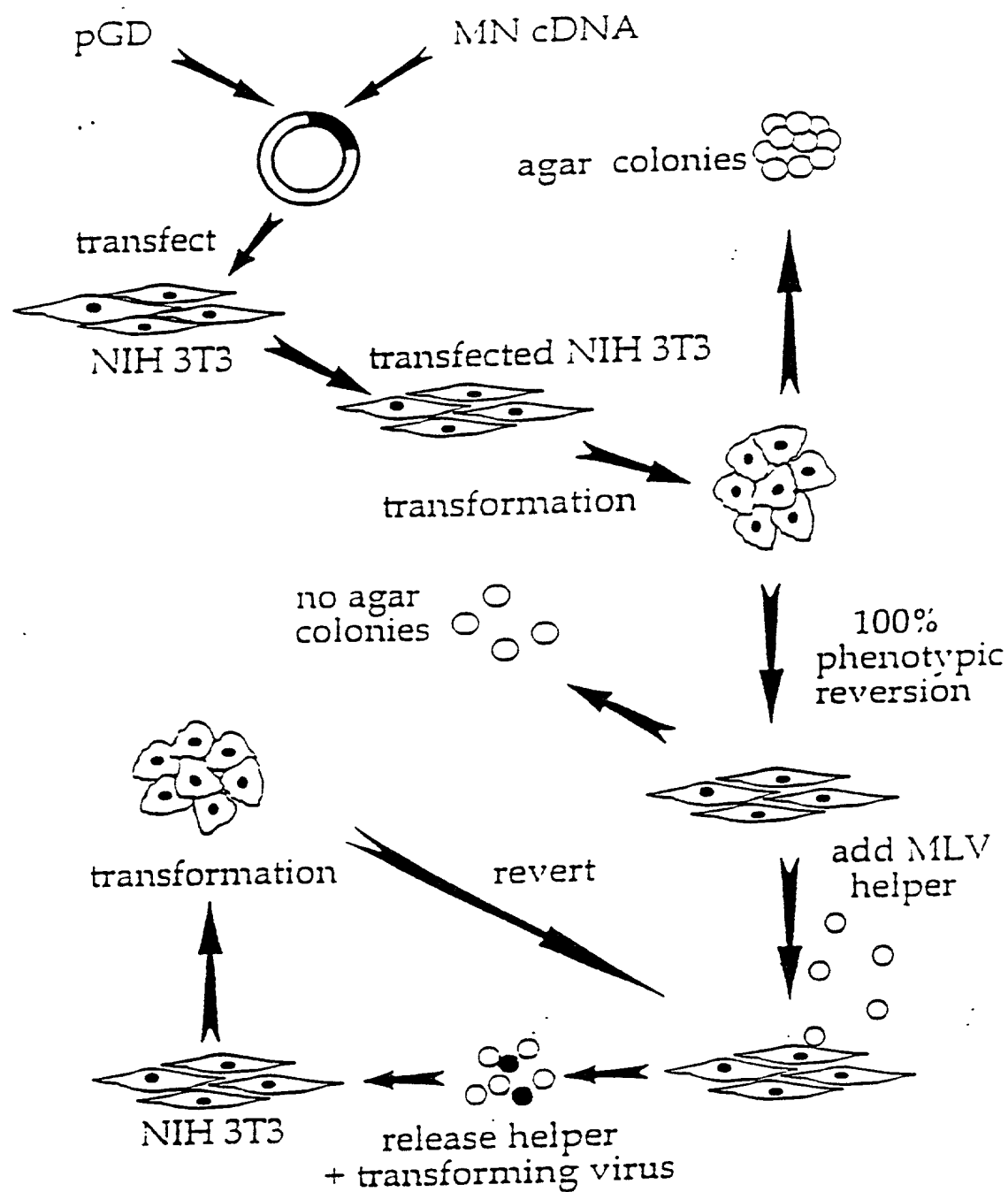


FIG.-8

**FIG._9**